## **AMENDMENTS TO THE CLAIMS:**

The following listing of claims will replace all prior versions and listings of claims in the specification.

## 1-27. (Cancelled)

- 28. (New) An isolated nucleic acid molecule comprising a sequence of nucleotides encoding or complementary to a sequence encoding a flavonoid 3',5' hydroxylase (F3'5'H), said nucleotide sequence selected from the list consisting of:
- (i) a nucleotide sequence encoding an amino acid sequence selected from SEQ ID NO:10 or SEQ ID NO: 12;
- (ii) a nucleotide sequence encoding an amino acid sequence having 90% similarity to an amino acid sequence selected from SEQ ID NO: 10 or SEQ ID NO: 12 after optimal alignment;
  - (iii) SEQ ID NO: 9 or SEQ ID NO: 11;
- (iv) a nucleotide sequence capable of hybridizing to a nucleotide sequence selected from SEQ ID NO:9 or SEQ ID NO: 11 or a complementary form thereof under high stringency conditions; and
- (v) a nucleotide sequence having 80% identity to a nucleotide sequence selected from SEQ ID NO: 9 or SEQ ID NO: 11.
- 29. (New) The isolated nucleic acid molecule of Claim 28 operably linked to a promoter.
- 30. (New) The isolated nucleic acid molecule of Claim 29 wherein the promoter is the CaMV35s promoter.
- 31. (New) A genetic construct comprising a nucleic acid molecule of any one of Claims 28 to 30.
- 32. (New) A genetically modified plant or progeny thereof comprising a nucleic acid molecule of Claim 28, wherein expression of said nucleic acid molecule results in an altered

flower color.

- 33. (New) The genetically modified plant of Claim 32 wherein said nucleic acid molecule is operably linked to a promoter.
- 34. (New) The genetically modified plant of Claim 33 wherein the promoter is the CaMV35s promoter.
- 35. (New) A genetically modified plant according to any one of Claims 32-34 wherein the plant is a rose or a progeny thereof.
- 36. (New) An isolated F3'5'H polypeptide comprising an amino acid sequence selected from the list consisting of:
  - (i) SEQ ID NO: 12;
  - (ii) sequences having at least 90% similarity to SEQ ID NO: 12;
  - (iii) SEQ ID NO: 10; and
  - (iv) sequences having at least 90% similarity to SEQ ID NO: 10.
- 37. (New) A fusion protein comprising at least one polypeptide according to Claim 36.